



## SEQUENCE LISTING

<110> Kalluri, Raghuram

<120> Anti-Angiogenic Proteins and Fragments and Methods of Use Thereof

<130> 02312/2085B (formerly 1440.1027-005)

<140> US 09/543,371

<141> 2000-04-04

<150> US 60/089,689

<151> 1998-06-17

<150> US 60/126,175

<151> 1999-03-25

<150> US 09/335,224

<151> 1999-06-17

<160> 18

<170> PatentIn version 3.1

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20 25 30	
ttg ctc tac gtg caa ggc aat gaa cgg gcc cat gga cag gac ttg ggc	144
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35 40 45	
acg gcc ggc agc tgc ctg cgc aag ttc agc aca atg ccc ttc ctg ttc	192
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50 55 60	
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Cys Asn Ile Asn Asn Val Cys Asn Phe Ala Ser Arg Asn Asp Tyr Ser	
65 70 75 80	
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Tyr Trp Leu Ser Thr Pro Glu Pro Met Pro Met Ser Met Ala Pro Ile	

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Pro Cys Pro Ser Gly Trp Ser Ser Leu Trp Ile Gly Tyr Ser Phe Val				
	130	135	140	
atg cac acc agc gct ggt gca gaa ggc tct ggc caa gcc ctg gcg tcc				480
Met His Thr Ser Ala Gly Ala Glu Gly Ser Gly Gln Ala Leu Ala Ser				
	145	150	155	160
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ctc gcc acc ata gag agg agc gag atg ttc aag aag cct acg ccg tcc				624
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	195	200	205	
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His Gly Arg Gly Thr Cys Asn Tyr Tyr Ala Asn Ala Tyr Ser Phe Trp  
180 185 190

Leu Ala Thr Ile Glu Arg Ser Glu Met Phe Lys Lys Pro Thr Pro Ser  
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 Asn Pro Gly Asp Val Cys Tyr Tyr Ala Ser Arg Asn Asp Lys Ser Tyr  
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 Trp Leu Ser Thr Thr Ala Pro Leu Pro Met Met Pro Val Ala Glu Asp  
 85 90 95  
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 Glu Ile Lys Pro Tyr Ile Ser Arg Cys Ser Val Cys Glu Ala Pro Ala  
 100 105 110  
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145	150	155	160
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Cys Leu Glu Asp Phe Arg Ala Thr Pro Phe Ile Glu Cys Asn Gly Gly			
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cgc ggc acc tgc cac tac tac gcc aac aag tac agc ttc tgg ctg acc			576
Arg Gly Thr Cys His Tyr Tyr Ala Asn Lys Tyr Ser Phe Trp Leu Thr			
	180	185	190
.acc att ccc gag cag agc ttc cag ggc tcg ccc tcc gcc gac acg ctc			624
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aag gcc ggc ctc atc cgc aca cac atc agc cgc tgc cag gtg tgc atg			672
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Arg Gly Thr Cys His Tyr Tyr Ala Asn Lys Tyr Ser Phe Trp Leu Thr  
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20 25 30

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Ser Cys Pro Glu Gly Thr Val Pro Leu Tyr Ser Gly Phe Ser Phe Leu
35 40 45

ttt gta caa gga aat caa cga gcc cac gga caa gac ctt gga act ctt 192
Phe Val Gln Gly Asn Gln Arg Ala His Gly Gln Asp Leu Gly Thr Leu
50 55 60

ggc agc tgc ctg cag cga ttt acc aca atg cca ttc tta ttc tgc aat 240
Gly Ser Cys Leu Gln Arg Phe Thr Thr Met Pro Phe Leu Phe Cys Asn
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Leu Ser Thr Pro Ala Leu Met Pro Met Asn Met Ala Pro Ile Thr Gly	
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aga gcc ctt gag cct tat ata agc aga tgc act gtt tgt gaa ggt cct	384
Arg Ala Leu Glu Pro Tyr Ile Ser Arg Cys Thr Val Cys Glu Gly Pro	
115 120 125	
gcg atc gcc ata gcc gtt cac agc caa acc act gac att cct cca tgt	432
Ala Ile Ala Ile Ala Val His Ser Gln Thr Thr Asp Ile Pro Pro Cys	
130 135 140	
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Pro His Gly Trp Ile Ser Leu Trp Lys Gly Phe Ser Phe Ile Met Phe	
145 150 155 160	
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165 170 175	
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Ser Cys Leu Glu Glu Phe Arg Ala Ser Pro Phe Leu Glu Cys His Gly	
180 185 190	
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Arg Gly Thr Cys Asn Tyr Tyr Ser Asn Ser Tyr Ser Phe Trp Leu Ala	
195 200 205	
tca tta aac cca gaa aga atg ttc aga aag cct att cca tca act gtg	672
Ser Leu Asn Pro Glu Arg Met Phe Arg Lys Pro Ile Pro Ser Thr Val	
210 215 220	
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Phe Val Gln Gly Asn Gln Arg Ala His Gly Gln Asp Leu Gly Thr Leu  
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Gly Ser Cys Leu Gln Arg Phe Thr Thr Met Pro Phe Leu Phe Cys Asn  
65 70 75 80

Val Asn Asp Val Cys Asn Phe Ala Ser Arg Asn Asp Tyr Ser Tyr Trp  
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Leu Ser Thr Pro Ala Leu Met Pro Met Asn Met Ala Pro Ile Thr Gly  
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Arg Ala Leu Glu Pro Tyr Ile Ser Arg Cys Thr Val Cys Glu Gly Pro  
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Ala Ile Ala Ile Ala Val His Ser Gln Thr Thr Asp Ile Pro Pro Cys  
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Pro His Gly Trp Ile Ser Leu Trp Lys Gly Phe Ser Phe Ile Met Phe  
145 150 155 160

Thr Ser Ala Gly Ser Glu Gly Thr Gly Gln Ala Leu Ala Ser Pro Gly  
165 170 175

Ser Cys Leu Glu Glu Phe Arg Ala Ser Pro Phe Leu Glu Cys His Gly  
180 185 190

Arg Gly Thr Cys Asn Tyr Tyr Ser Asn Ser Tyr Ser Phe Trp Leu Ala  
195 200 205

Ser Leu Asn Pro Glu Arg Met Phe Arg Lys Pro Ile Pro Ser Thr Val  
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